Problem 3 (6 Points)

SciKit-Learn only offers a few built-in preprocessors, such as MinMax and Standard scaling. However, it also offers the ability to create custom data transformation functions, which can be integrated into your pipeline. In this problem, you will implement a log transform and observe how using it changes a regression result.

Start by running this cell to import modules and load data:

```
In [2]: import numpy as np
        import matplotlib.pyplot as plt
        from sklearn.model_selection import train_test_split
        from sklearn.metrics import mean_squared_error
        from sklearn.pipeline import Pipeline
         from sklearn.preprocessing import FunctionTransformer
        from sklearn.svm import SVR
        def plot(X_train, X_test, y_train, y_test, model=None,log = False):
             plt.figure(figsize=(5,5),dpi=200)
             if model is not None:
                X_fit = np.linspace(min(X_train)-0.15, max(X_train)+0.2)
                 y_fit = model.predict(X_fit)
                 plt.plot(np.log(X_fit+1) if log else X_fit, y_fit,c="red",label="Prediction")
             if log:
                X_train = np.log(X_train+1)
                 X_{\text{test}} = \text{np.log}(X_{\text{test+1}})
             plt.scatter(X_train,y_train,s=30,c="powderblue",edgecolors="navy",linewidths=.5,12
             plt.scatter(X_test,y_test,s=30,c="orange",edgecolors="red",linewidths=.5,label="Texts.")
             plt.legend()
             plt.xlabel("log(x+1)" if log else "x")
             plt.ylabel("y")
             plt.show()
        x = np.array([5.83603919, 1.49205924, 2.66109578, 9.40172515, 6.47247125, 0.3763]
        X = x.reshape(-1,1)
        y = np.array([4.32538472e+00, -5.59312420e+00, -4.57455876e+00, 4.23667057e+01, 1.04]
        X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=1, train_size=@)
```

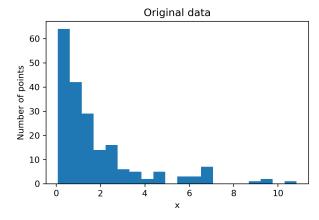
Distribution of x data

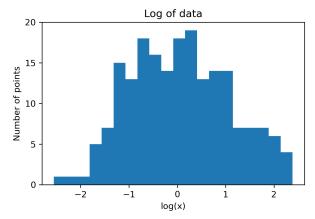
Let's visualize how the original input feature is distributed, alongside the log of the data -notice that performing this log transformation makes the data much closer to normally
distributed.

```
In [3]: plt.figure(figsize=(12,3.4),dpi=300)
   plt.subplot(1,2,1)
   plt.hist(x,bins=20)
   plt.xlabel("x")
   plt.ylabel("Number of points")
```

```
plt.title("Original data")

plt.subplot(1,2,2)
plt.hist(np.log(x),bins=20)
plt.xlabel("log(x)")
plt.ylabel("Number of points")
plt.title("Log of data")
plt.ylim(0,20)
plt.yticks([0,5,10,15,20])
plt.show()
```

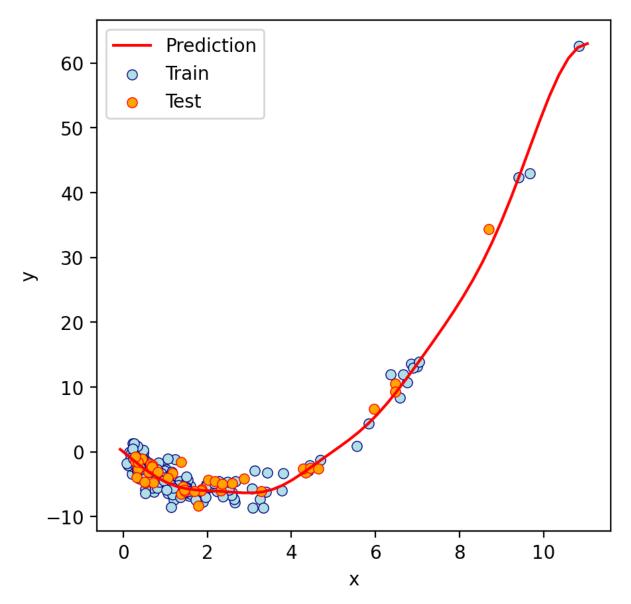




No log transform

First, we do support vector regression on the untransformed inputs. The code to do this has been provided below.

Training MSE: 2.071006155233616 Testing MSE: 1.9453578716771704



With log transform

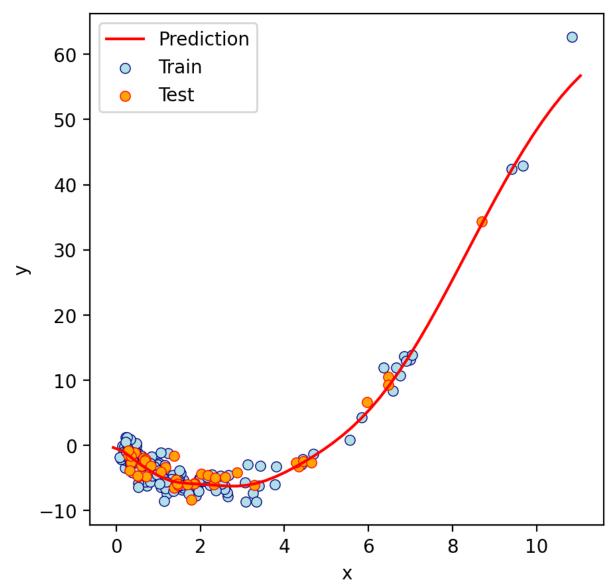
Notice that the data are not spread uniformly across the x axis. Instead, most input data points have low values -- this is a roughly "log normal" distribution. If we take the log of the input, we saw it was more normally distributed, which can improve machine learning model results in some cases. The transform function has been given below. Add this to a new pipeline, train the pipeline, and compute the train MSE and test MSE. Show a plot as above. Note the subtle change in behavior of the fitting curve.

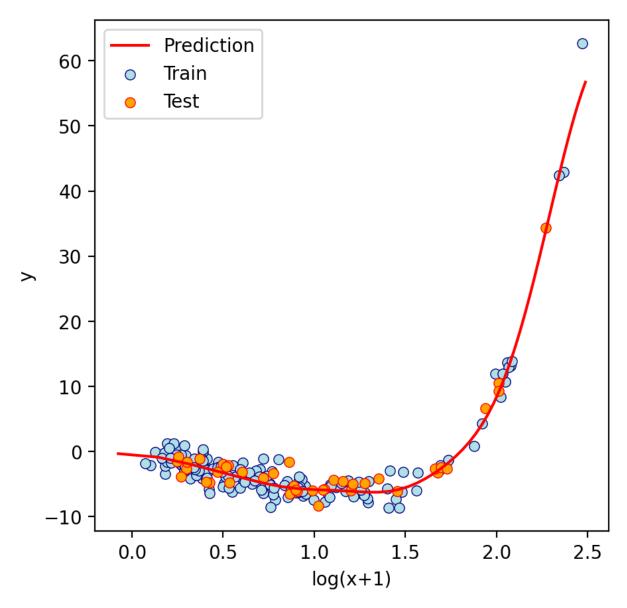
Also, make another plot setting the log argument to True. This will show the scaling of the x-axis used by the model.

```
In [5]: def log_transform(x):
    return np.log(x + 1.)

transform = FunctionTransformer(log_transform)
model = SVR(C=100)
```

Training MSE: 2.3139214731606486 Testing MSE: 1.7200875366333022





In []: